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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/715,418

DATE: 10/19/2001
TIME: 11:25:00

Input Set : A:\09-715,418 sequence listing Attorney docket 10716-12.txt
Output Set: N:\CRF3\10192001\I715418.raw

3 <110> APPLICANT: CURAGEN CORPORATION
4 RASTELLI, LUKA
6 <120> TITLE OF INVENTION: WNT-REGULATED CYTOKINE-LIKE POLYPEPTIDE AND NUCLEIC
7 ACIDS ENCODING SAME
9 <130> FILE REFERENCE: 10716/12
11 <140> CURRENT APPLICATION NUMBER: 09/715,418
C--> 12 <141> CURRENT FILING DATE: 2000-11-19
14 <150> PRIOR APPLICATION NUMBER: 60/166,177
15 <151> PRIOR FILING DATE: 1999-11-18
17 <160> NUMBER OF SEQ ID NOS: 48
19 <170> SOFTWARE: PatentIn Ver. 2.1
21 <210> SEQ ID NO: 1
22 <211> LENGTH: 212
23 <212> TYPE: DNA
24 <213> ORGANISM: Mus sp.
26 <400> SEQUENCE: 1
27 gaattcagtg atgtagagag ggccattgag acactcatca agaacttcca taaatactct 60
28 gtggcgggta aaaaggaaac actgacccct gctgagcttc gagacctggt tacccagcag 120
29 ctgccacacc tcatgccgag caactgtggg ttagaagaga aaattgccaa cctgggcaac 180
30 tgtaatgact cgaaactgga gtttggaagc tt 212
33 <210> SEQ ID NO: 2
34 <211> LENGTH: 1670
35 <212> TYPE: DNA
36 <213> ORGANISM: Mus sp.
38 <220> FEATURE:
39 <221> NAME/KEY: modified_base
40 <222> LOCATION: (1541)
41 <223> OTHER INFORMATION: "n" represents a, t, c, g, other or unknown
43 <400> SEQUENCE: 2
44 tcaggtgagc tggctcctcc atcctgtctc ccagctgcca gcaggtctcc ccctcctcta 60
45 ggtagatcat gatccatcag ctcctgtggg gcaggctata ggacagacga caaaactcaa 120
46 ctcacagaag gaaggaccag tgtaccagga acgatgggac agtgtcggtc agccaatgct 180
47 gaggatgccc aagaattcag tgatgtagag agggccattg agacactcat caagaacttc 240
48 cataaatact ctgtggcggg taaaaaggaa acactgaccc ctgctgagct tcgagacctg 300
49 gttacccagc agctgccaca cctcatgccg agcaactgtg ggttagaaga gaaaattgcc 360
50 aacctgggca actgtaatga ctcgaaactg gagtttggaa gcttctggga gttgattgga 420
51 gaagcagcca agagtgtgaa gatggagagg cctgttactc ggagctgagg acttctactt 480
52 ggaacttgtt gggggtgttg gggatagggg agttttagag gcactggaaa taaaaccctc 540
53 aatgcccacc acccccttcc ccagcctgca cctctcctca ttgctgcaat gttcacgttc 600
54 aggacaggct tccctgtggg ctccatggag ctcctgggtc cagaagtcct catctcaagg 660
55 gagctcaggg ggtgggttgg ggctggagag gatatgcagg gatcctggaa gggtaagggc 720
56 caagcaattt ggtagtaggg gaagggcaga aaggaactgg gttatggaag tgatccaaag 780
57 agcagggatg ggaatctggc tgcatatttg gtcctgaaaa gggtgtctga gaacctaccc 840
58 ccttctaatc ttgtcccacc taaactgtag ttgtctgccc tgtgctatcc ttgctgcttc 900
59 cagctctgcc ccatcctcct tccagtgtct gttcctgagt aggggcaggg gaaataggag 960
60 cagagttgca aaagaggctg aggagggcat gacttcatca ctttggggtg agaggaccag 1020
61 ctagatgctt gggcatttat ggtagttatt ttatatcatt tgattaataa aaatattgga 1080

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    62 aaatgtaaag aaaaaaaaag aaaaaaacat ggggccgaaa ccttatcccc cttgagtagg 1140
    63 gtgatatttt gcgtgtgcaa tgggcggcct gttttcgaga ggcggtgaca tggggaaaac 1200
    64 atgggggtgt accaaacctt aaccgccttt taggggaaac acccctttg ccgcaagtgg 1260
    65 gttaataacg gaagaagccc ggccggattg cccttcacaa gagtctcccg cggtagatgc 1320
    66 ggatgggaca gccccccttcg gcggcgttta gagcggcgtg tgtgtggttt ctacgcgaat 1380
    67 agggataaat attgtggcgg cgccgaggga gtgtgtgtgt tgcgcgcctg cttctgtgga 1440
    68 ggtggtgtgt cccaaaaact aaaagggccc ttttgtgcgc gttagtttgc tctagcagag 1500
W--> 69 tccgctgcac atattttggt gggcgtgtcc gtgccgcccg nggtggtgct tgttgctggc 1560
    70 gtggcgtggg gtgggtgtgg ttgcgggggt ggtcgtgttg ggtgtgtgcg tgcgcgcggg 1620
    71 ggccgtgtgt gtgtgtggtt gcatgataag gttagagtga gtgagagcgg            1670
    74 <210> SEQ ID NO: 3
    75 <211> LENGTH: 131
    76 <212> TYPE: PRT
    77 <213> ORGANISM: Mus sp.
    79 <400> SEQUENCE: 3
    80 Ser Ile Ser Ser Cys Gly Ala Gly Tyr Arg Thr Asp Asp Lys Thr Gln
    81   1               5                  10                  15
    83 Leu Thr Glu Gly Arg Thr Ser Val Pro Gly Thr Met Gly Gln Cys Arg
    84              20                  25                  30
    86 Ser Ala Asn Ala Glu Asp Ala Gln Glu Phe Ser Asp Val Glu Arg Ala
    87          35                  40                  45
    89 Ile Glu Thr Leu Ile Lys Asn Phe His Lys Tyr Ser Val Ala Gly Lys
    90      50                  55                  60
    92 Lys Glu Thr Leu Thr Pro Ala Glu Leu Arg Asp Leu Val Thr Gln Gln
    93  65                  70                  75                  80
    95 Leu Pro His Leu Met Pro Ser Asn Cys Gly Leu Glu Glu Lys Ile Ala
    96                  85                  90                  95
    98 Asn Leu Gly Asn Cys Asn Asp Ser Lys Leu Glu Phe Gly Ser Phe Trp
    99             100                 105                 110
   101 Glu Leu Ile Gly Glu Ala Ala Lys Ser Val Lys Met Glu Arg Pro Val
   102         115                 120                 125
   104 Thr Arg Ser
   105     130
   108 <210> SEQ ID NO: 4
   109 <211> LENGTH: 357
   110 <212> TYPE: DNA
   111 <213> ORGANISM: Homo sapiens
   113 <220> FEATURE:
   114 <221> NAME/KEY: modified_base
   115 <222> LOCATION: (231)
   116 <223> OTHER INFORMATION: "n" represents a, t, c, g, other or unknown
   118 <220> FEATURE:
   119 <221> NAME/KEY: modified_base
   120 <222> LOCATION: (337)
   121 <223> OTHER INFORMATION: "n" represents a, t, c, g, other or unknown
   123 <400> SEQUENCE: 4
   124 ataggacaac agaactctca ccaaaggacc agacacagtg agcaccatgg gacagtgtcg 60
   125 gtcagccaac gcagaggatg ctcaggaatt cagtgatgtg gagagggcca ttgagaccct 120
   126 catcaagaac tttcaccagt actccgtgga gggtgggaag gagacgctga ccccttctga 180
```

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W--> 127 gctacgggac ctggtcaccc agcagctgcc ccatctcatg ccgagcaact ntggcctgga 240
128 agagaaaatt gccaacctgg gcagctgcaa tgactctaaa ctggagttca ggagtttctg 300
W--> 129 ggagctgatt ggagaagcgg ccaagagtgt gaagctngag aggactgtcc gggggca 357
132 <210> SEQ ID NO: 5
133 <211> LENGTH: 379
134 <212> TYPE: DNA
135 <213> ORGANISM: Homo sapiens
137 <400> SEQUENCE: 5
138 gaattccaga gggagttctc agtgcccccg gacaggcctc tccagcttca cactcttggc 60
139 cgcttctcca atcagctccc agaaactcct gaactccagt ttagagtcat tgcagctgcc 120
140 caggttggca attttctctt ccaggccaca gttgctcggc atgagatggg gcagctgctg 180
141 ggtgaccagg tcccgtagct cagaaggggt cagcgtctcc ttcccaccct ccacggagta 240
142 ctggtgaaag ttcttgatga gggtctcaat ggccctctcc acatcactga attcctgagc 300
143 atcctctgcg ttggctgacc gacactgtcc catggtgctc actgtgtctg gtcctttggt 360
144 gagagttctg ttgtcctat 379
147 <210> SEQ ID NO: 6
148 <211> LENGTH: 118
149 <212> TYPE: PRT
150 <213> ORGANISM: Homo sapiens
152 <400> SEQUENCE: 6
153 Asp Asn Arg Thr Leu Thr Lys Gly Pro Asp Thr Val Ser Thr Met Gly
154 1 5 10 15
156 Gln Cys Arg Ser Ala Asn Ala Glu Asp Ala Gln Glu Phe Ser Asp Val
157 20 25 30
159 Glu Arg Ala Ile Glu Thr Leu Ile Lys Asn Phe His Gln Tyr Ser Val
160 35 40 45
162 Glu Gly Gly Lys Glu Thr Leu Thr Pro Ser Glu Leu Arg Asp Leu Val
163 50 55 60
165 Thr Gln Gln Leu Pro His Leu Met Pro Ser Asn Cys Gly Leu Glu Glu
166 65 70 75 80
168 Lys Ile Ala Asn Leu Gly Ser Cys Asn Asp Ser Lys Leu Glu Phe Arg
169 85 90 95
171 Ser Phe Trp Glu Leu Ile Gly Glu Ala Ala Lys Ser Val Lys Leu Glu
172 100 105 110
174 Arg Pro Val Arg Gly His
175 115
178 <210> SEQ ID NO: 7
179 <211> LENGTH: 20
180 <212> TYPE: DNA
181 <213> ORGANISM: Artificial Sequence
183 <220> FEATURE:
184 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
186 <400> SEQUENCE: 7
187 cttgatgagg gtctcaatgg 20
190 <210> SEQ ID NO: 8
191 <211> LENGTH: 26
192 <212> TYPE: DNA
193 <213> ORGANISM: Artificial Sequence
195 <220> FEATURE:

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196 <223> OTHER INFORMATION: Description of Artificial Sequence: Probe
198 <400> SEQUENCE: 8
199 ccacatcact gaattcctga gcatcc 26
202 <210> SEQ ID NO: 9
203 <211> LENGTH: 20
204 <212> TYPE: DNA
205 <213> ORGANISM: Artificial Sequence
207 <220> FEATURE:
208 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
210 <400> SEQUENCE: 9
211 cagacacagt gagcaccatg 20
214 <210> SEQ ID NO: 10
215 <211> LENGTH: 98
216 <212> TYPE: PRT
217 <213> ORGANISM: Homo sapiens
219 <400> SEQUENCE: 10

```
Met Ala Ala Glu Pro Leu Thr Glu Leu Glu Glu Ser Ile Glu Thr Val
 1               5                  10                  15
Val Thr Thr Phe Phe Thr Phe Ala Arg Gln Glu Gly Arg Lys Asp Ser
            20                  25                  30
Leu Ser Val Asn Glu Phe Lys Glu Leu Val Thr Gln Gln Leu Pro His
        35                  40                  45
Leu Leu Lys Asp Val Gly Ser Leu Asp Glu Lys Met Lys Ser Leu Asp
    50                  55                  60
Val Asn Gln Asp Ser Glu Leu Lys Phe Asn Glu Tyr Trp Arg Leu Ile
65                  70                  75                  80
Gly Glu Leu Ala Lys Glu Ile Arg Lys Lys Lys Asp Leu Lys Ile Arg
                85                  90                  95
Lys Lys
```

242 <210> SEQ ID NO: 11
243 <211> LENGTH: 110
244 <212> TYPE: PRT
245 <213> ORGANISM: Homo sapiens
247 <400> SEQUENCE: 11

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Met Ser Gln Leu Glu Arg Asn Ile Glu Thr Ile Ile Asn Thr Phe His
 1               5                  10                  15
Gln Tyr Ser Val Lys Leu Gly His Pro Asp Thr Leu Asn Gln Gly Glu
            20                  25                  30
Phe Lys Glu Leu Val Arg Lys Asp Leu Gln Asn Phe Leu Lys Lys Glu
        35                  40                  45
Asn Lys Asn Glu Lys Val Ile Glu His Ile Met Glu Asp Leu Asp Thr
    50                  55                  60
Asn Ala Asp Lys Gln Leu Ser Phe Glu Glu Phe Ile Met Leu Met Ala
65                  70                  75                  80
Arg Leu Thr Trp Ala Ser His Glu Lys Met His Glu Gly Asp Glu Gly
                85                  90                  95
Pro Gly His His His Lys Pro Gly Leu Gly Glu Gly Thr Pro
            100                 105                 110
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270 <210> SEQ ID NO: 12

RAW SEQUENCE LISTING DATE: 10/19/2001
PATENT APPLICATION: US/09/715,418 TIME: 11:25:00

Input Set : A:\09-715,418 sequence listing Attorney docket 10716-12.txt
Output Set: N:\CRF3\10192001\I715418.raw

271 <211> LENGTH: 37
272 <212> TYPE: PRT
273 <213> ORGANISM: Unknown Organism
275 <220> FEATURE:
276 <223> OTHER INFORMATION: Description of Unknown Organism: 3-100/ICaBP type
277 calcium binding protein
279 <400> SEQUENCE: 12
280 Ser Asn Cys Gly Leu Glu Glu Lys Ile Ala Asn Leu Gly Ser Cys Asn
281 1 5 10 15
283 Asp Ser Lys Leu Glu Phe Arg Ser Phe Trp Glu Leu Ile Gly Glu Ala
284 20 25 30
286 Ala Lys Ser Val Lys
287 35
290 <210> SEQ ID NO: 13
291 <211> LENGTH: 37
292 <212> TYPE: PRT
293 <213> ORGANISM: Unknown Organism
295 <220> FEATURE:
296 <223> OTHER INFORMATION: Description of Unknown Organism: 3-100/ICaBP type
297 calcium binding protein
299 <400> SEQUENCE: 13
300 Asp Val Glu Arg Ala Ile Glu Thr Leu Ile Lys Asn Phe His Gln Tyr
301 1 5 10 15
303 Ser Val Glu Gly Gly Lys Glu Thr Leu Thr Pro Ser Glu Leu Arg Asp
304 20 25 30
306 Leu Val Thr Gln Gln
307 35
310 <210> SEQ ID NO: 14
311 <211> LENGTH: 19
312 <212> TYPE: PRT
313 <213> ORGANISM: Unknown Organism
315 <220> FEATURE:
316 <223> OTHER INFORMATION: Description of Unknown Organism: Bacterial type II
317 secretion system protein F
319 <400> SEQUENCE: 14
320 Val Thr Gln Gln Leu Pro His Leu Met Pro Ser Asn Cys Gly Leu Glu
321 1 5 10 15
323 Glu Lys Ile
327 <210> SEQ ID NO: 15
328 <211> LENGTH: 10
329 <212> TYPE: PRT
330 <213> ORGANISM: Unknown Organism
332 <220> FEATURE:
333 <223> OTHER INFORMATION: Description of Unknown Organism: Ubiquitin
334 carboxyl-terminal hydrolases family
336 <400> SEQUENCE: 15
337 Ala Asn Ala Glu Asp Ala Gln Glu Phe Ser
338 1 5 10
341 <210> SEQ ID NO: 16

VERIFICATION SUMMARY DATE: 10/19/2001
PATENT APPLICATION: **US/09/715,418** TIME: 11:25:01

Input Set : **A:\09-715,418 sequence listing Attorney docket 10716-12.txt**
Output Set: **N:\CRF3\10192001\I715418.raw**

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:69 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:127 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:129 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4